

Figur 1

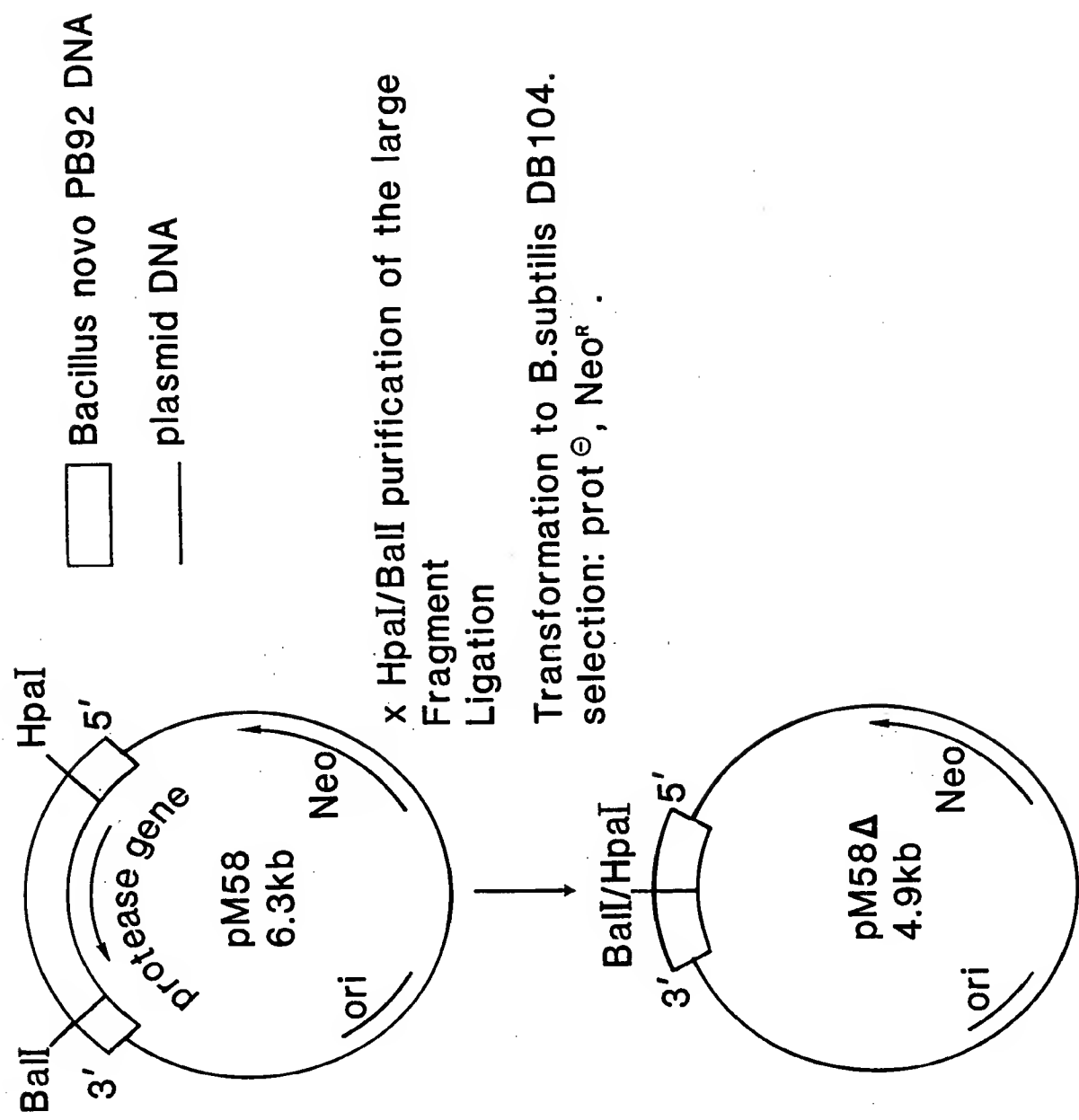


Figure 2

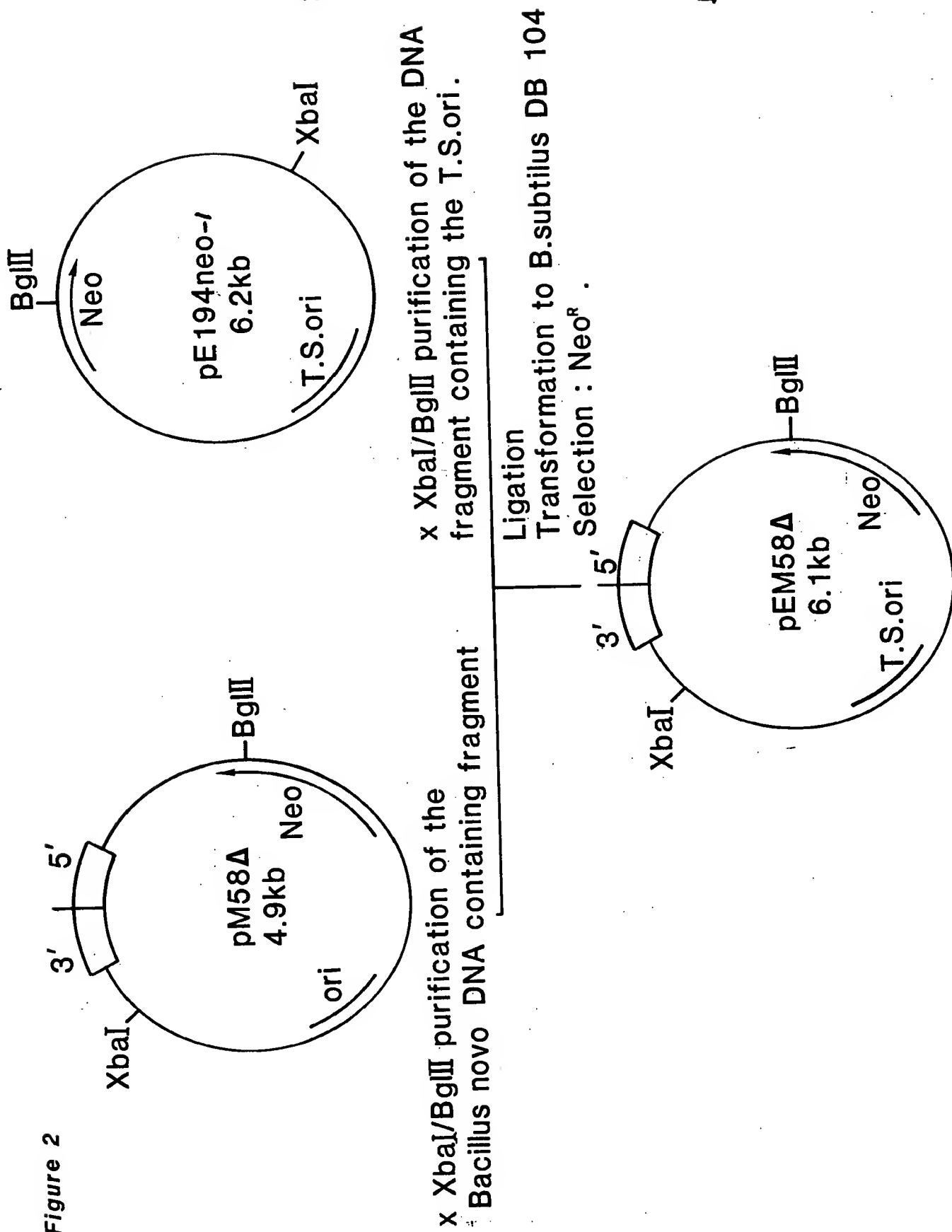


Figure 3

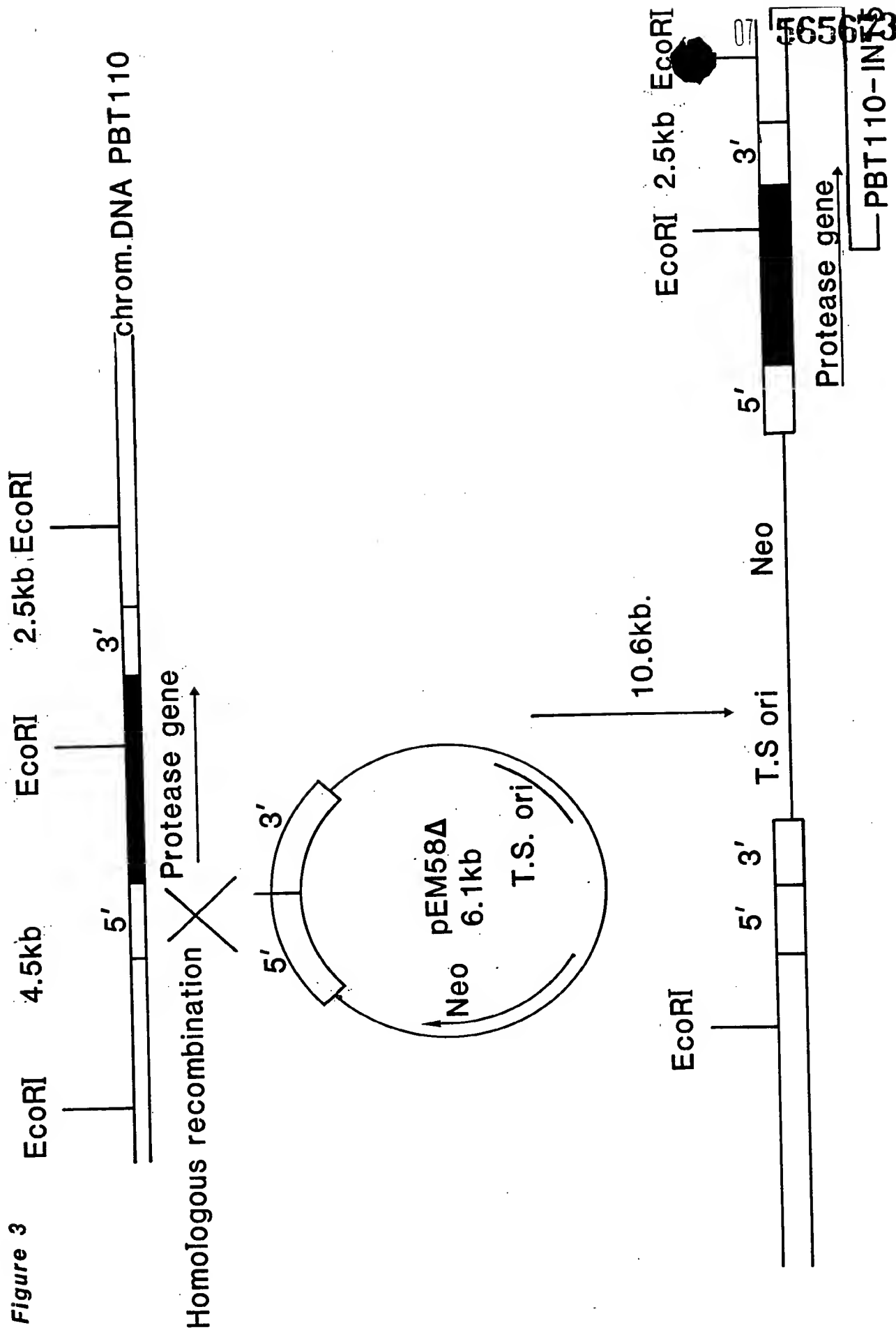
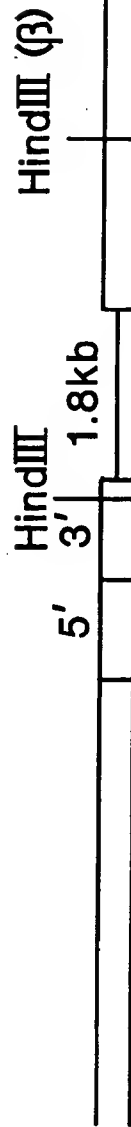


Figure 4



07 565623



HindIII . . . . . 40 . . . . . 07 565673 80  
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 . . . . . 120 . . . . . 160  
 AAAATATTGCTTATGCAACAGTTCATAATGATGAGCAAACCCCTCACATGCATTTAGGTGTTGTGCCTATGCGTGATGGA  
 . . . . . 200 . . . . . 240  
 AAACCTGCAAGGAAAAAATGTGTTTAAATCGTCAAGAACTGTTATGGCTACAAGATAAATTCCTCCGAGCATATGAAAAACA  
 . . . . . 280 . . . . . 320  
 GGGTTTTGAGTTGAAGCGTGGTGAACGTGGCTCTGACCGTAAACATATTGAGACAGCTAAATTTAAAAAACAACTTTGG  
 . . . . . XbaI . . . . . 360 . . . . . 400  
 AAAAAGAGATTGATTTTCTAGAAAATGAATACGGTAAGGAAAATATGCTGTATGCGACTGTCCATCTGGATGAAAGAGTC  
 . . . . . 440 . . . . . 480  
 CCACATATGCACCTTTGGTTTTGTCCCTTTAACAGAGGACGGGAGATTGTCTGCAAAAGAACAGTTAGGCAACAAGAAAGA  
 . . . . . 520 . . . . . 560  
 CTTTACTCAATTACAAGATAGATTTAATGAGTATGTGAATGAGAAAGGTTATGAACTTGAAAGAGGCACGTCCAAAGAGG  
 . . . . . 600 . . . . . 640  
 TTACAGAACGAGAACATAAAGCGATGGATCAGTACAAGAAAGATACTGTATTTTCATAAACAGGAACTGCAAGAAGTTAAG  
 . . . . . 680 . . . . . 720  
 GATGAGTTACAGAAGGCAAATAAGCAGTTACAGAGTGGAATAGAGCATATGAGGTCTACGAAACCCTTTGATTATGAAAA  
 . . . . . 760 . . . . . 800  
 TGAGCGTACAGGTTTGTCTCTGGACGTGAAGAGACTGGTAGAAAGATATTAAGTCTGATGAATTTGAACGCCTGCAAG  
 . . . . . PstI . . . . . 840 . . . . . 880  
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 . . . . . 920 . . . . . 960  
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 . . . . . 1000 . . . . . 1040  
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 . . . . . 1080 . . . . . 1120  
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 . . . . . 1160 . . . . . 1200  
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 . . . . . 1240 . . . . . 1280  
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 . . . . . 1320 . . . . . 1360  
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 . . . . . 1400 . . . . . 1440  
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 . . . . . plasm.seq. chrom.seq. . . . . 1520  
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 . . . . . 1560 . . . . . 1600  
 AATTTCTAGGCAACAACAACGCCGCTATTCTTGACGGTGGGATTTCAGGCTTGAAGAACAGTGGCTACCCGACAACGGGA  
 . . . . . HindIII  
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figure 5A

figure 5.B

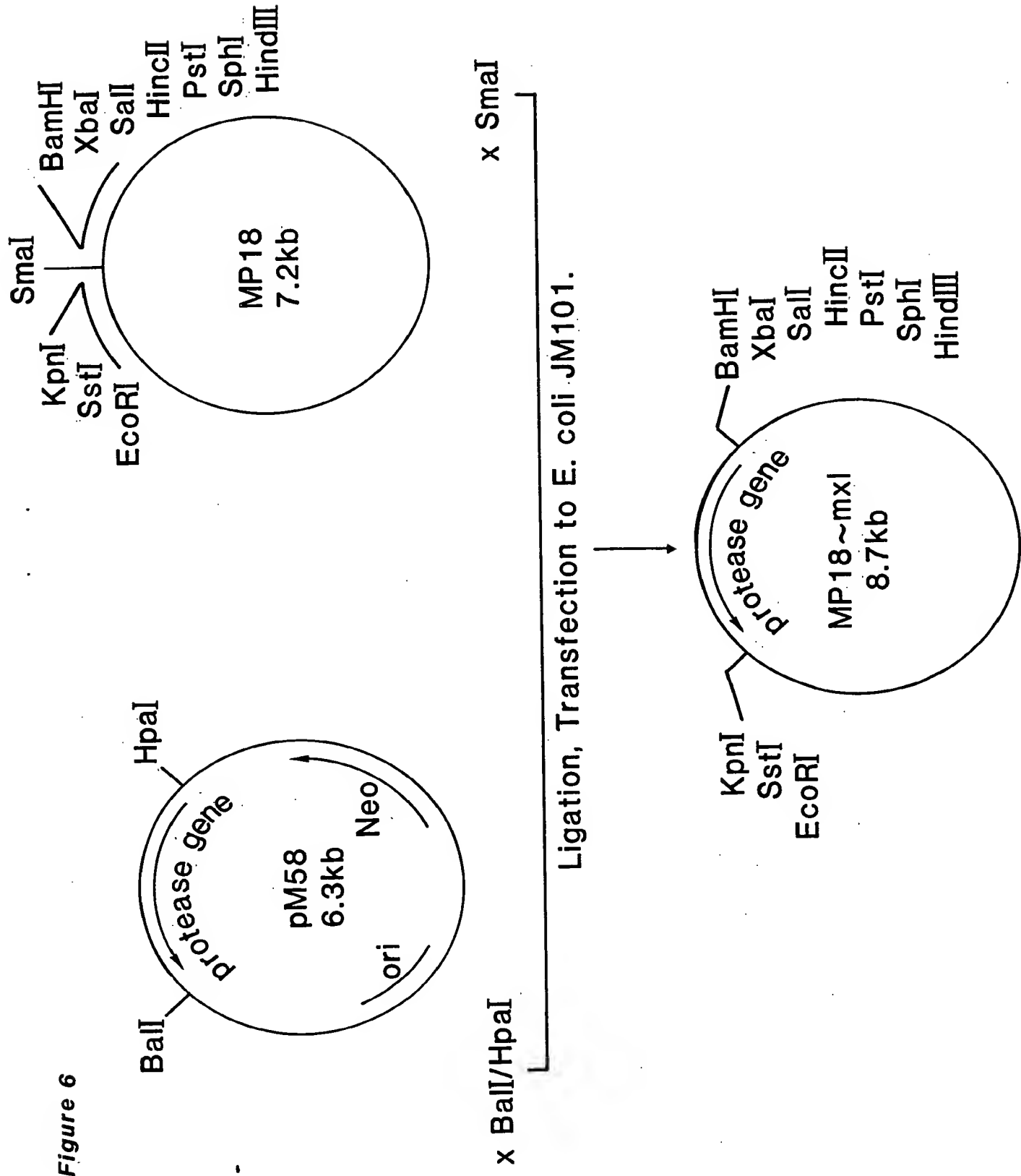


Figure 6

## Sequence of pBHA-1

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                                150                                180
GAAAGCAAGC TGATAAACCG ATACAATTAA AGGCTCCTTT TGGAGCCTTT TTTTTTGGAG
                                210                                240
ATTTTCAACG TGAAAAAATT ATTATTCGCA ATTCCAAGCT CTGCCTCGCG CGTTTCGGTG
                                270                                300
ATGACGGTGA AAACCTCTGA CACATGCAGC TCCCGGAGAC GGTCACAGCT TGTCTGTAAG
                                330                                360
CGGATGCAGA TCACGCGCCC TGTAGCGGCG CATTAAAGCGC GGCGGGTGTG GTGGTTACGC
                                390                                420
GCAGCGTGAC CGCTACACTT GCCAGCGCCC TAGCGCCCGC TCCTTTCGCT TTCTTCCCCT
                                450                                480
CCTTTCTCGC CACGTTCGCC GGCTTTCCCC GTCAAGCTCT AAATCGGGGG CTCCCTTTAG
                                510                                540
GGTTCCGATT TAGTGCTTTA CGGCACCTCG ACCCCAAAAA ACTTGATTAG GGTGATGGTT
                                570                                600
CACGTAGTGG GCCATCGCCC TGATAGACGG TTTTTCGCCC TTTGACGTTG GAGTCCACGT
                                630                                660
TCTTTAATAG TGGACTCTTG TTCCAAACTG GAACAACACT CAACCCTATC TCGGTCTATT
                                690                                720
CTTTTGATTT ATAAGGGATT TTGCCGATTT CGGCCTATTG GTTAAAAAAT GAGCTGATTT
                                750                                780
AACAAAAATT TAACGCGAAT TTTAACAAAA TATTAACGTT TACAATTTGA TCTGCGCTCG
                                810                                840
GTCGTTCCGC TGC GCGAGC GGTATCAGCT CACTCAAAGG CGGTAATACG GTTATCCACA
                                870                                900
GAATCAGGGG ATAACGCAGG AAAGAACATG TGAGCAAAAG GCCAGCAAAA GGCCAGGAAC
                                930                                960
CGTAAAAAGG CCGCGTTGCT GGC GTTTTTTC CATAGGCTCC GCCCCCTGA CGAGCATCAC
                                990                                1020
AAAAATCGAC GCTCAAGTCA GAGGTGGCGA AATCCGACAG GACTATAAAG ATACCAGGCG
                                1050                                1080
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```

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1170 1200  
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1230 1260  
CCCGACCGCT GCGCCTTATC CGGTAACAT CGTCTTGAGT CCAACCCGGT AAGACACGAC

1290 1320  
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1350 1380  
GCTACAGAGT TCTTGAAGTG GTGGCCTAAC TACGGCTACA CTAGAAGGAC AGTATTTGGT

1410 1440  
ATCTGCGCTC TGCTGAAGCC AGTTACCTTC GGAAAAAGAG TTGGTAGCTC TTGATCCGGC

1470 1500  
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1530 1560  
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1650 1680  
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1710 1740  
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1770 1800  
TCCATAGTTG CCTGACTCCC CGTCGTGTAG ATAACCTACGA TACGGGAGGG CTTACCATCT

1830 1860  
GGCCCCAGTG CTGCAATGAT ACCGCGAGAC CCACGCTCAC CGGCTCCAGA TTTATCAGCA

1890 1920  
ATAAACCAGC CAGCCGGAAG GGCCGAGCGC AGAAGTGGTC CTGCAACTTT ATCCGCCTCC

1950 1980  
ATCCAGTCTA TTAATTGTTG CCGGGAAGCT AGAGTAAGTA GTTCGCCAGT TAATAGTTTG

2010 2040  
CGCAACGTTG TTGCCATTGC TGCAGGCATC GTGGTGTCAC GCTCGTCGTT TGGTATGGCT

2070 2100  
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2130 2160  
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2190 2220  
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2250 2280  
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7170 7200  
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Figure 8

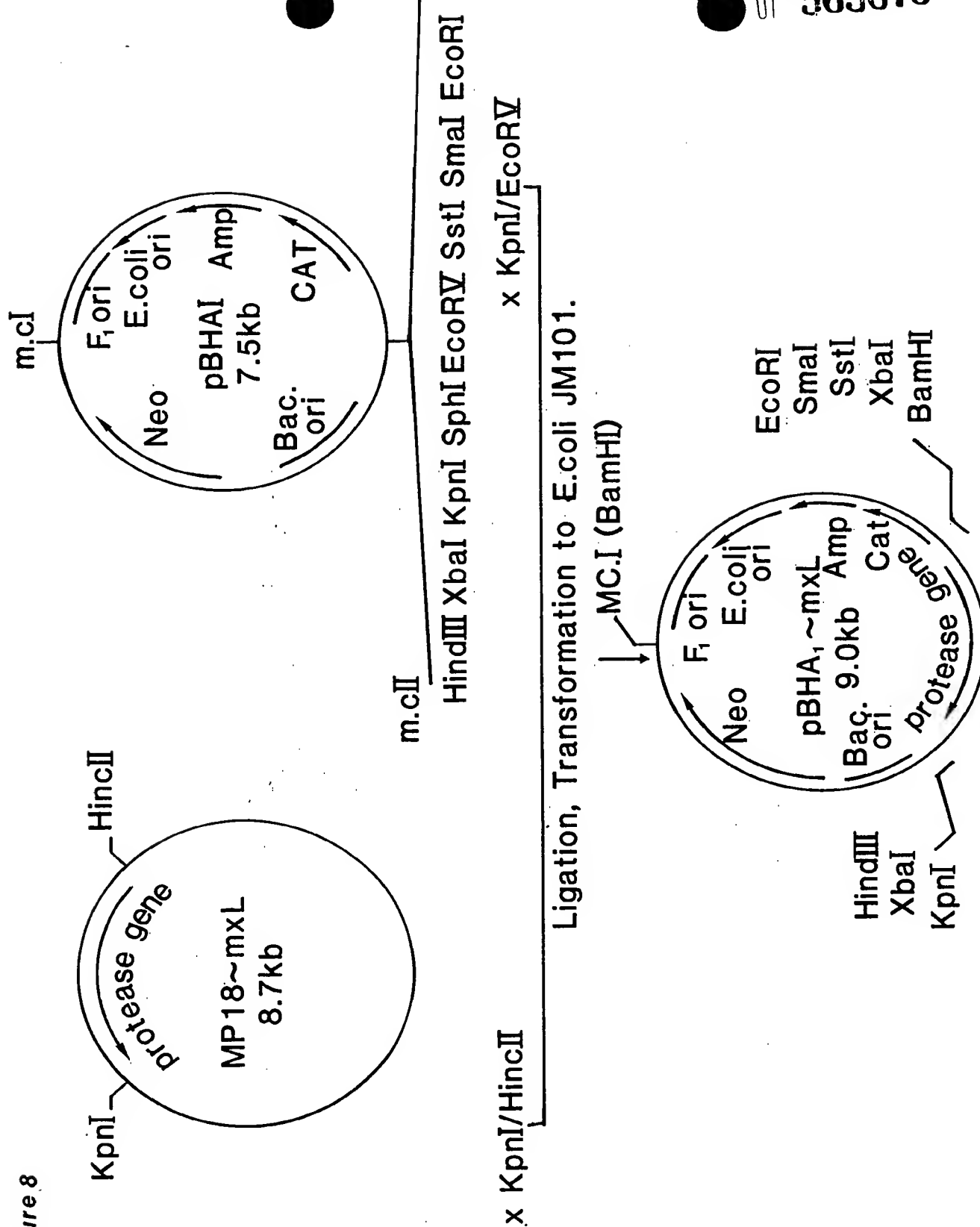


Figure 9

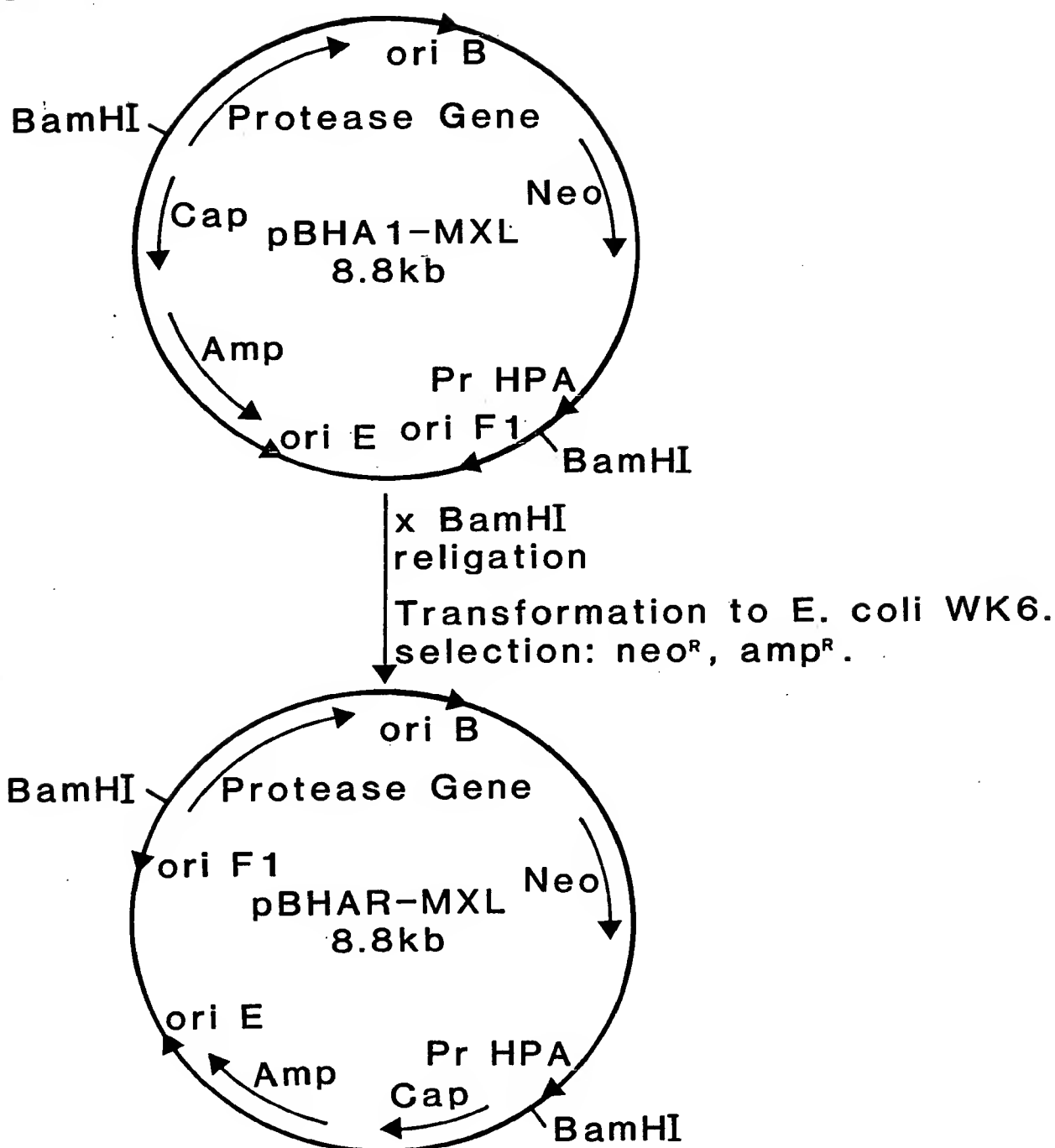
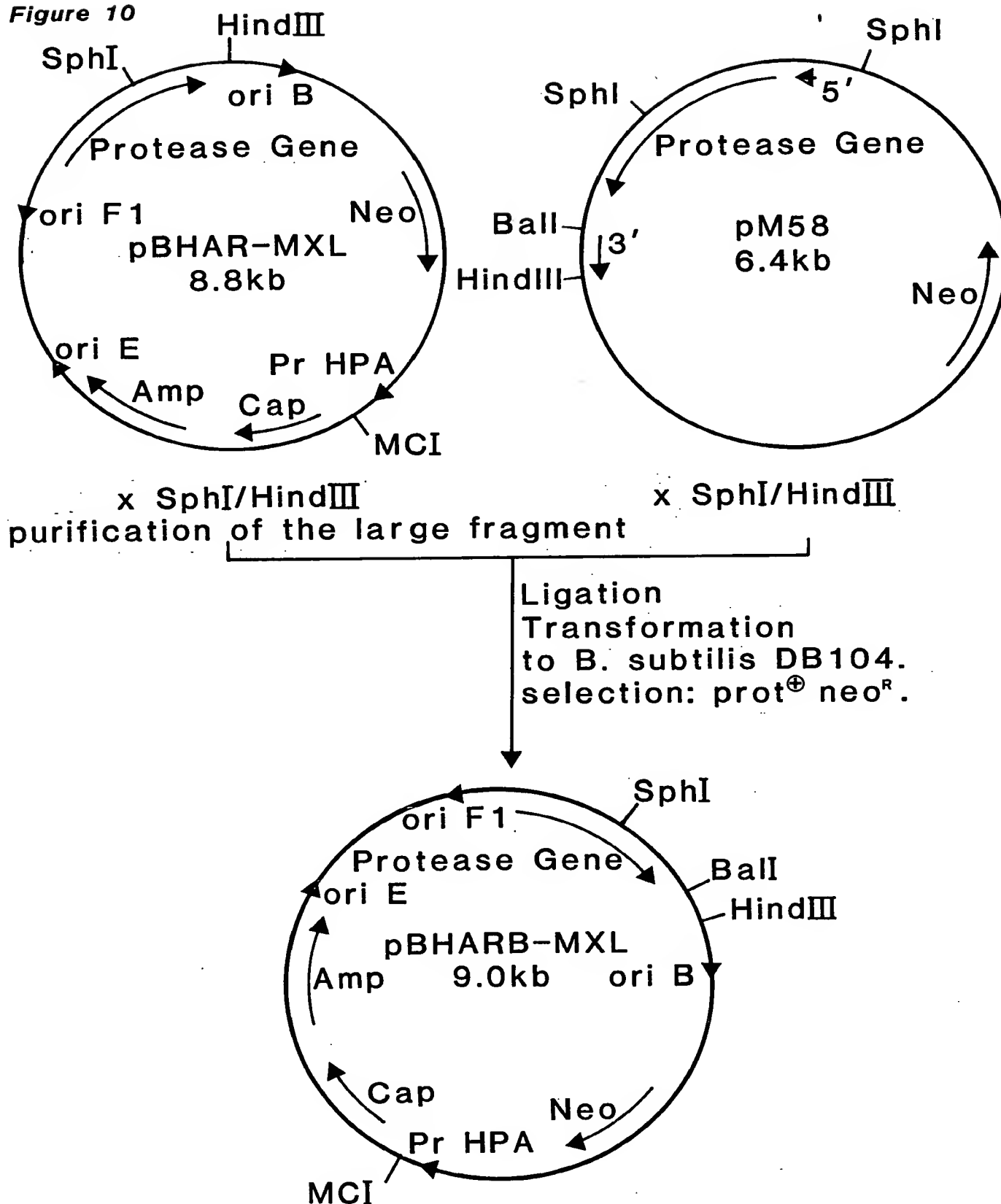
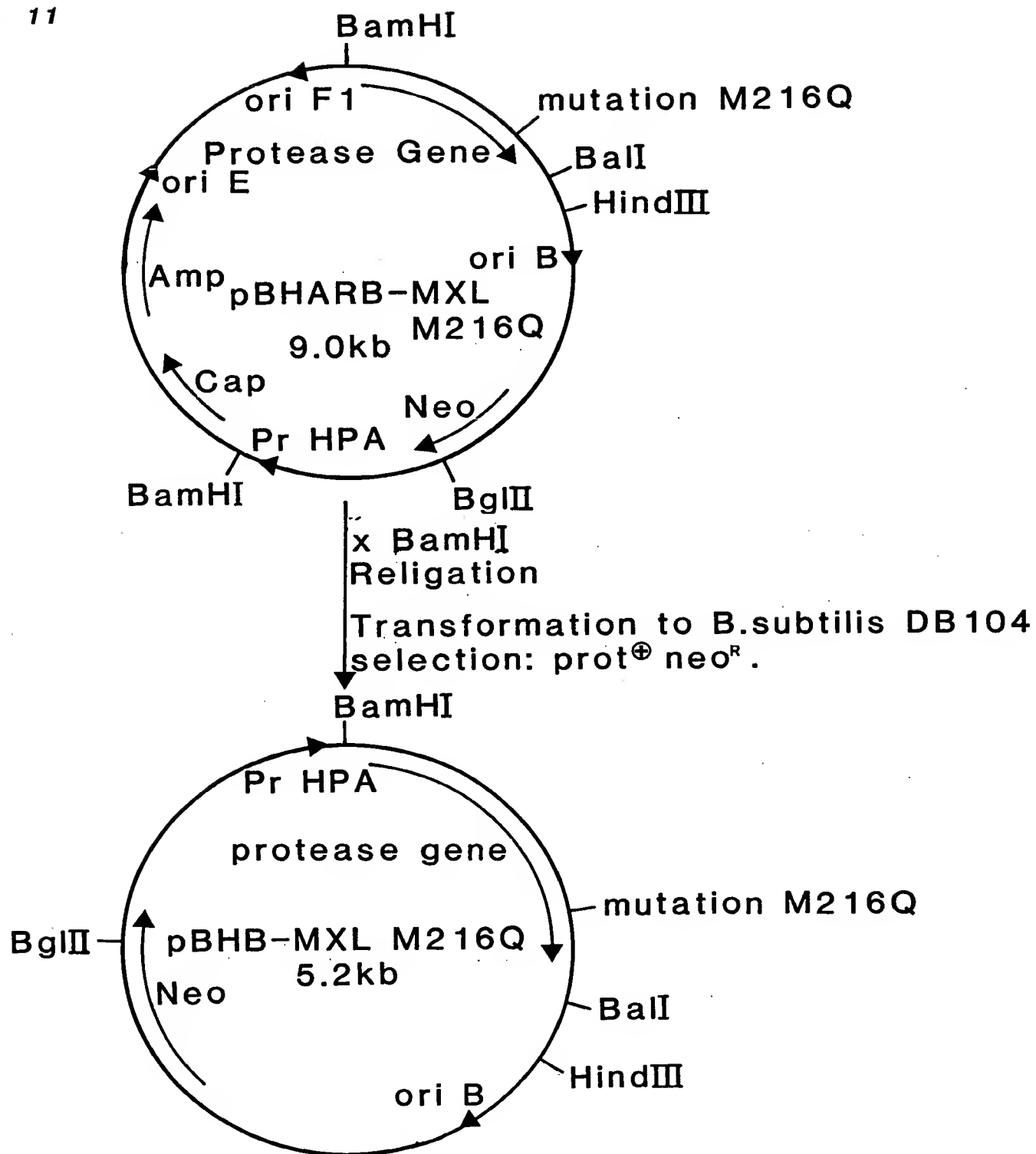


Figure 10



Figur 11



07 565673

Figure 12

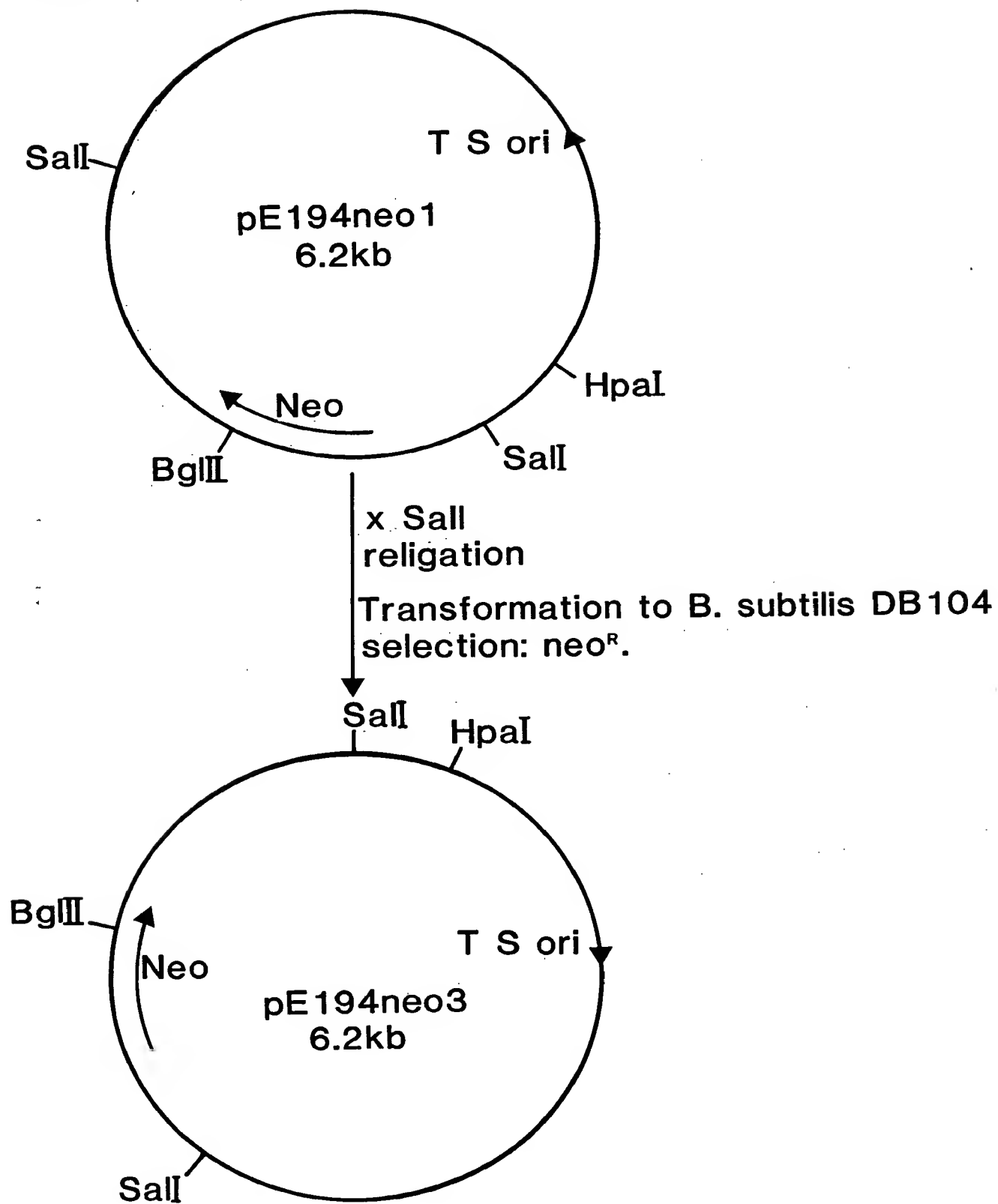


Figure 13

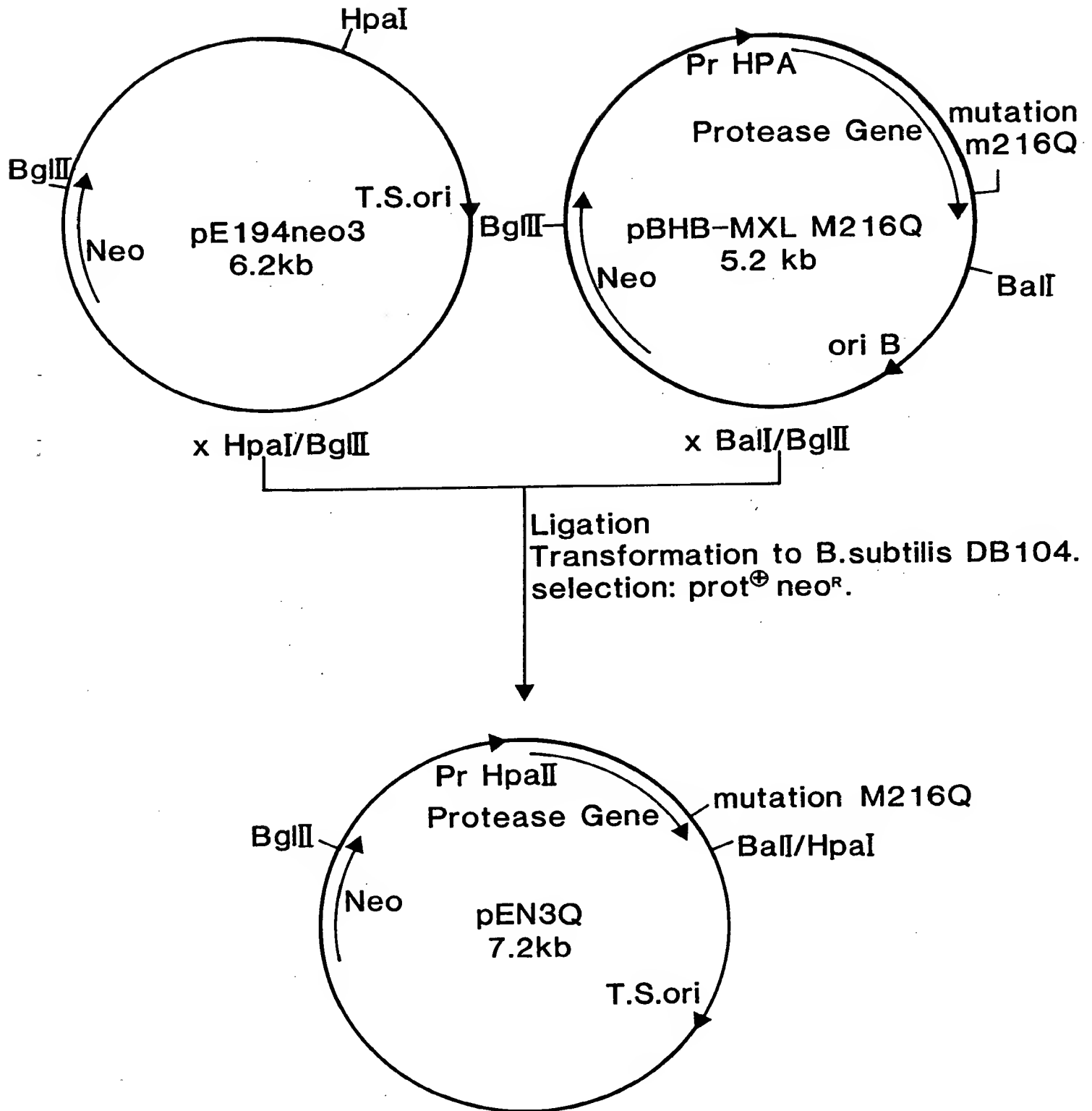


Figure 14

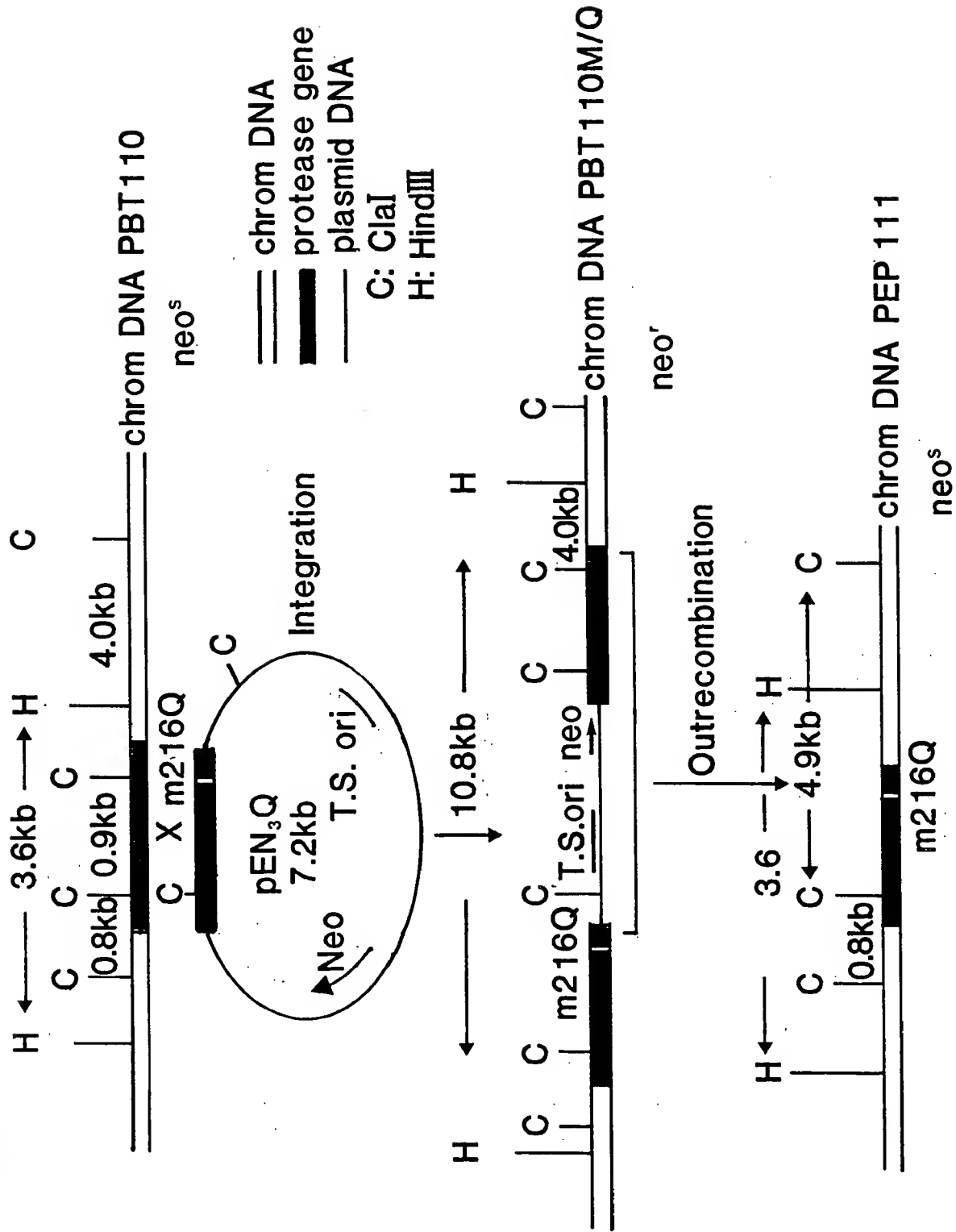


Figure 15

